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Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2008; month=10; day=30; hr=15; min=50; sec=20; ms=833;
]

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Application No: 10583877

Version No: 2.0

Input Set:

Output Set:

Started: 2008-10-01 14:55:38.061

Finished: 2008-10-01 14:55:40.208

Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 147 ms

Total Warnings: 44

Total Errors: 2

No. of SeqIDs Defined: 97

Actual SeqID Count: 97

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (51)
W 213	Artificial or Unknown found in <213> in SEQ ID (52)
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W 213	Artificial or Unknown found in <213> in SEQ ID (54)
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W 213	Artificial or Unknown found in <213> in SEQ ID (57)
W 213	Artificial or Unknown found in <213> in SEQ ID (58)
W 213	Artificial or Unknown found in <213> in SEQ ID (59)
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W 213	Artificial or Unknown found in <213> in SEQ ID (69)
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Input Set:

Output Set:

Started: 2008-10-01 14:55:38.061
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Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 147 ms
Total Warnings: 44
Total Errors: 2
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Error code	Error Description
	This error has occurred more than 20 times, will not be displayed
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (87)

SEQUENCE LISTING

<110> Ellis, Jonathan H

Eon-Duval, Alexandre

Grundy, Robert I

Hussain, Farhana

McAdam, Ruth

Plumpton, Christopher

Prinjha, Rabinder K

Wison, Paul A

<120> NOGO - A NEUTRALISING IMMUNOGLOBULINS FOR TREATMENT OF NEUROLOGICAL DISEASES

<130> PB60608

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<141> 2006-06-21

<150> PCT/GB2004/005325

<151> 2004-12-20

<150> GB 0329684.5

<151> 2003-12-22

<150> GB 0329711.6

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<400> 37

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Trp Met His Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
35 40 45

Gly Asn Ile Asn Pro Ser Asn Gly Gly Thr Asn Tyr Asn Glu Lys Phe
50 55 60

Lys Ser Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr
65 70 75 80

Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys
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Glu Leu Gly Gln Gly Tyr Trp Gly Gln Gly Thr Thr Leu Thr Val Ser
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<210> 38

<211> 115

<212> PRT

<213> Mus musculus

<400> 38

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Ala Met Ser Trp Val Arg Gln Thr Pro Glu Lys Arg Leu Glu Trp Val
35 40 45

Ala Ser Ile Ser Asp Gly Gly Ser Tyr Thr Tyr Tyr Pro Asp Asn Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Asn Leu Tyr
65 70 75 80

Leu Gln Met Ser His Leu Lys Ser Glu Asp Thr Ala Met Tyr Tyr Cys
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Ala Lys Glu Leu Leu Phe Asp Tyr Trp Gly Gln Gly Thr Thr Leu Thr
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Val Ser Ser
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<210> 39

<211> 113

<212> PRT

<213> Mus musculus

<400> 39

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Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ala Phe Ser Ser Tyr
20 25 30

Trp Met Asn Trp Val Lys Gln Arg Pro Gly Lys Gly Leu Glu Trp Ile
35 40 45

Gly Gln Ile Tyr Pro Gly Asp Gly Asp Thr Asn Tyr Asn Gly Lys Phe
50 55 60

Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala Tyr
65 70 75 80

Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Phe Cys
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Ala Val Arg Phe Asp Tyr Trp Gly Gln Gly Thr Thr Leu Thr Val Ser

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110

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<210> 40

<211> 112

<212> PRT

<213> Mus musculus

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20 25 30

Asp Gly Lys Thr Tyr Leu Asn Trp Phe Leu Gln Arg Pro Gly Gln Ser
35 40 45

Pro Gln Leu Leu Ile Tyr Leu Met Ser Thr Arg Ala Ser Gly Val Ser
50 55 60

Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Glu Ile
65 70 75 80

Ser Arg Val Lys Ala Glu Asp Val Gly Val Tyr Tyr Cys Gln Gln Leu
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Val Glu Tyr Pro Leu Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys
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<210> 41

<211> 112

<212> PRT

<213> Mus musculus

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20 25 30

Asn Gly Asn Thr Tyr Leu His Trp Tyr Leu Gln Lys Pro Gly Gln Ser
35 40 45

Pro Lys Leu Leu Ile Tyr Lys Val Ser Asn Arg Phe Ser Gly Val Pro
50 55 60

Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile
65 70 75 80

Ser Arg Val Glu Ala Glu Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser
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<210> 42

<211> 112

<212> PRT

<213> Mus musculus

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20 25 30

Asn Gly Asn Thr Tyr Leu Tyr Trp Phe Leu Gln Arg Pro Gly Gln Ser
35 40 45

Pro Gln Leu Leu Ile Tyr Arg Met Ser Asn Leu Ala Ser Gly Val Pro
50 55 60

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			85						90					95	

Leu	Glu	Tyr	Pro	Leu	Thr	Phe	Gly	Ala	Gly	Thr	Lys	Leu	Glu	Leu	Lys
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<400> 44

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cctggaaagg gtcttgagtg gattggacag atttactctg gagatggtga tactaactac 180

aacggaaagt tcaagggcaa ggccacactg actgcagaca aatcctccag cacagcctac 240

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<212> DNA

<213> Mus musculus

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